OIPE



ENTERED



TECH CENTER 1600/2900 DATE: 05/05/200

TIME: 07:36:18

RAW SEQUENCE LISTING

1 - 110 APPLICANT: Widner, William

PATENT APPLICATION: US/09/834,271A

Input Set: N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\I834271A.raw

```
Sloma, Alan
              Thomas, Michael E.
      4 <120 - TITLE OF INVENTION: Methods For Producing A polypeptide In a
             Bacillus Cell
      • <130 FILE REFERENCE: 5455.200-US
      " -:140 - CUERENT APPLICATION NUMBER: 09/834,271A
     M (141 - CURLENT FILING DATE: 2001-04-12
      → HISS - FRIOR APPLICATION NUMBER: US/04/258,377
     1 | <151. Phior Filing DATE: 1999-02-26
     11 <15 / PELOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/031,442
W--> 12 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-26
     13 4160 NUMBER OF SEQ ID NOS: 35
     14 <170 SOFTWARE: FastSEQ for Windows Version 3.0
     16 .210 SEP ID NO: I
     1) GM12. TYPE: DNA
19 GM130 GERANISM: Bacillus
     20 HAGIN SEQUENCE: 1
             sattggrett aagggeeegg gaegteaage ttategatge ggateegegg eege
                                                                                       54
     25 kd110 seq 18 No: 2
     24 -03110 FENGTH: 51
     원 HELER TYPE: DNA
     26 CHES CHEMANISM: Bacillus
     27 -4000 SEQUENCE: 2
              rongaattire egggeeetge agttegaata getaegeeta ggegeeggeg e
                                                                                       51
     3 4.110 SHQ II NO: 3
     M 40110 DEKGTH: B8
     N. KUISH IYPE: DNA
     by K.113: CECANISM: Bacillus
     34 HACO SEQUENCE: 3
             agotangeet taagggood ggacgtogag otdaagottg oggoegocat ggtogadg
     :" -1.10: SEQ ID NO: 4
     % -1.11 BENGTH: 58
     30 C. 120 TYPE: DNA.
     10 -0.13: OFCANISM: Bacillus
     90 - 4000 SEQUENCE: 4
              torqqaafto cogggoootg cagotogagt togaacgoog goggtaccag ofgottaa
     44 C.10: SEQ ID NO: 5
     45 -CHIEF LENGTH: 37
    46 CHIES TYPE: DNA
    47 KE13: CFGANISM: Bacillus
     48 <400; SEQUENCE: 5
```

RAW SEQUENCE LISTING DATE: 08/05/2003 PATENT APPLICATION: US/09/834,271A TIME: C7:36:18

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\1834271A.raw

49		ctccgggcsc atctgagete	tataaaaatg	aggaggg	37
51	$< \cap 1 \cap +$	SEO ID NO: 6			
5.2	·1.:1: ·	IENGTH: 17			
53	·1.:1. ·	IYEE: DUA			
54	1.11 h	OF GANISM: Facillus			
ţ5 °,	$\{ \cdot \} \cap \{ \cdot \}$	FEQUENCE: ϵ			
E 1 + 1		iotogratuo atacacaaaa	aaacgct		27
Ē, -	410 -	SEC ID NO: "			
£, .		LEDGTH: 87			
6.	·1.11. ·	TYLE: DNA			
61	4.11 - 4	(F. AMUSM: Pacillus			
Ġ.	1.4	SM, UEDCE: 7			
6 -		reaggactta agggccgcat	gegteettet	ttgtgct	37
		SEM: 10 NO: 8			
6,		IENGTH: ED			
6.	·1.:1 ·	TYPE: DNA			
(al-	41.1	CE WANTEM: Famillus			
6,11	454:11 ×	SEQUENCE: 8			
		r againtro titleaatgtg	taacatatga		30
		SE, ID NO: 9			
		LEIGTH: 48			
		TYPE: ONA			
j t	·: ::	OF ANTEM: Facillus			
		SEQUEDOE: 9			
		triggelita agggeotgea	atcgattgtt	tgagaaaaga ag	42
· , •		3F, 10 No: 10			
差子	-1.711	LF:MGTH: 48			
		TYPE: DMA			
		OF WANTEM: Facillus			
7	-(41)	SEQUENCE: 10			
÷ .;		trigadones attitottat	acaaattata	ttttacatat cag	43
$\in G$		Sho ID No: 11			
Ė		LFTGTH: 30			
		TYFE: DNA			
814	-0.37 5"-	OF PANISM: Pacillus			
		SPOURDON: 11			
		gammacecogy agetttcagt	gaagtacgtg		30
		SB. ID NO: 12			
		LF1.FTH: 19			
		TYFE: DNA			
		OFGANISM: Pacillus			
		SEQUENCE: 12			
94.		quipiegtiae aattcaaag			19
1(4)		·Šký ID NO: 13			
		· LENGTH: 27			
		TYFE: DNA			
		· CRGANISM: Bacillus			
		· SEQUENCE: 13			
105		gggccctcga aacgtaagat	gaaacct		27
			100		

RAW SEQUENCE LISTING

DATE: 08/05/2003 TIME: 07:36:18 PATENT APPLICATION: US/09/834,271A

Imput Set : N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\1834271A.raw

		SED ID NO: 14					
		LENGTH: 29					
		TYFE: DNA					
		OPMANIEM: Bacillus					
		SEQUENCE: 14	7.1				200
1	. 5 1 .	priotizata atabataatt	ttcaaactg				29
		SE, TO NO: 15 LENGTH: 31					
		ITTE: DNA					
		OFGANUSM: Bacillus SEQUENCE: 15					
		-chyomien: 15 -chyocatoso áttgtgaaat					21
		SE, ID No: 16					1
		LENGTH: 23					
		TYPE: DNA					
		OFGANISM: Bacillus					
		SEQUENCE: 16					
		paintentato istaatiaag	ctt				23
		2F, 10 NO: 17	500				
		DETARTH: 25					
		TYPE: SOLA					
		OF FAMILFM: Bacillus					
		SEQUENCE: 13					
		quietoquac figitoatgt	gaa				23
		SPG TO No: 1:	-				
1 15	1.1	tengra: 13					
1 . 7	1.	TYPE: FNA					
1	1.1	GFGANISM: Babillus					
1:P	$\cdot,\cdot,0\cdots$	SEQUENCE: 18					
1 : i :		gayetesisa tacataatit	tca				23
14.	\cdot 1:	SEQ ID NO: 19					
		DECLETH: 44					
		TYFE: INA					
		OFFRANCSN: Bacillus					
	()	SEQUENCE: 19					
1 -		gnaafaang qqggttgaca	ttattttact	gatatgtatā	atat		4.4
		SEQ 20 NO: 30					
		LFNGTH: 48					
		TYFE: DNA					
		OFGANISM: Bacillus					
	- [-] () (1) -	SEQUENCE: 20			4		1.0
15.5		astaabatga ctatacatat	tatattaaac	atattetttt	acctcgag		48
		SEQ 10 NO: 21					
		LENGTH: 3050 TYPE: CNA					
		CFGANISM: a:Bacillus					
		SEQUENCE: 21					
161	~ + U U - '	togaaacgta agatgaaacc	ttamataaan	atacttttt	tattacsett	daadaattat	60
162		taatgttaag ottaattaaa					120
ے رہ ہ		camegerady credatedada	gacaacaccc	ergantegra	a c g c c c c c c c	adageadydd	4 4. 0

RAW SEQUENCE LISTING

DATE: 08/05/2003 PATENT APPLICATION: US/09/834,271A TIME: 07:36:19

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\I834271A.raw

163					cttcttcaga		120
164					aatgabatab		5:1
15°					gtgaattate		3:10
1 600			_		caagtagaaa		56)()
1 ()	ottgatagos	ttactatacc	taacatgatg	tagtattasa	tgaatatgta	aatatattta	47
16.5					tattggaatg		480
16	caatagaata	gtgtataaat	tatttatott	gaaaggaggg	atgodtaaaa	acgaagaaca	L
1 7	ttaaaaaacat	atatttgcac	ogtotaatgg	atttatgaaa	aatdattta	toagtttgaa	600
171	aattatgtat	tatgataaga	aagggaggaa	gaaaaatgaa	toogaacaat	cgaagtgaac	E) E) (_1
1 /	atgatabaat	aaaaactact	gaaaataatg	aggt godaad	taaccatgtt	caatatoott	7. 0
1	tagoggaaac	tocaaatoca	acactagaag	atttaaatta	taaagagtti	ttaagaatga	$^{2} \in \mathcal{O}$
17.4	otycagataa	taatacggaa	gcactagata	gototadaad	aaaagatgto	attossassg	5.440
1 70	gbatttdbgt	agtaggtgat	atastaggag	tagtaggttt	coogtitiggt	ggagogottg	эÇО
1	tittjogittita	tabaaacttt	ttaaatabta	tttggodaag	tgaagacccg	tggaaggett	46731
1	ttatggaaca	agtagaagca	ttgatggato	agaaaatago	tgattatgda	aaaaataaag	1:0.50
1	otottgbaga	gttacagggc	cttcaaaata	atglogaaga	statgsgagt	gcattgagtt	1:05:1
1 , .	catggcáaaa	aaatootgtg	agttcacqaa	atopapatag	coaggggogg	ataagagaga	114.1
1 : .:	tgiltitatak	agcagaaagt	cattttogta	attbaatgob	ttögittgöä	atitictggat	1,. (): (
1 1	abgaggttbt	atttotaaca	adatatgcad	aagdugddaa	cacacattta	titttadtas	1. 41.1
1	aagaogotba	aatttatgga	gaagaatggģ	gatacqaaaa	agaagatatt	gotgaatttt	1
1 :	ataasagaca	actaaaactt	acgcaagaat	atactgacca	ttgtgtdaaa	tggtataatg	1 580
1 ;					ggtaaacttt		1.;.4:1
1 🖓 🖰	gcagagagat	gacattaaca	gtattagatt	taattgcact	atttocattg	tatgatgttd	1 (11)
180	ggotatacco	aaaagaagtt	aaaaccgaat	taacaagaga	ogttitaaca	gatecasttg	15.610
15%					taatatagaa		1
1 %					toababgbgg		1
18 1	gatattatgg	aaatqactot	ttdaattatt	ggtooggtaa	ttatgtttca	actagaccaa	$1 \sim 10$
1 90	-				aaataaatoo		1800
1 %:	tabaaaattt	agaatttaat	ggagasasag	totatagago	ogtagoaaat	adaastottg	1 10
1 3.	aggtatggod	gtoogotgtá	tattdaggtg	ttadaaaagt	ggaatttagd	caatataatg	1
1 90	atcaaacaga	tgaagbaagt	acacaaacgt	acgacticaaa	aagaaatgtt	ggbgbgggtba	1 4.50
194	gotgggatto	tatogatõaa	ttgodtodag	aaadaadaga	tgaatotota	gaaaagggat	2 40
1.3					gggtagtaga		21-1
1 300	cagtattaac	ttggadadat	aaaagtgtag	acttttttaa	catgattgat	togaaaaaaa	33.66
197	ttäpäpääjät	toogttagta	aaggcatata	agttacaato	tiggtigettes	gttgtbgcag	:
19%					aaatggaagt		1100
19 (agotagaatt		3. 40
200	_				accatttaat		1.10
201					ttdätttaat		24.00
20:					aggogtbaba		2500
20.5					agtgaattaa		11. 11.
,) (h. ;					ataaaaaaat		11 41
201					gtagatgaag		.1700
200					taaagggaac		The fall
.:01			_		tagecagaaa		.: N: 1- Q
.:08					tagetteaag		11880
.1011					tatgtcccat		1000
.310					cataagtatg		3000
211			aaaagccctc			3	3050
	- 1-255500			,, c c c c	-)		

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/834,271A TIME: 07:36:19

DATE: 08/05/2003

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt
Gutput Set: N:\CRF4\08052003\I834271A.raw

```
213 42105 SET ID No: 22
314 -011 - DENGTH: 17
215 COLE TYPE: EMA
216 C.13 · OF CANLEM: Pacillus
31 - 40 - SEQUENCE: 13
                                                                                     17
.:1
          igarettalig godtgca
225 (216 - SEQ TO NO: 23
221 -011 - DEDGTH: .2
dd. -0.4. - TYFE: DWA
22 × 4.17 * ORGANISM: Fabillus
BB4 - 0400 SEQUENCE: MB
                                                                                      22
With thouse monstituted to
327 K. 10 - SEO ID NO: 74
dar -1.11 - LEDSTH: . :
AND BUTLE TYPE: DNA
Large 1 to 05 JANISM: Babillus
NAT - 440: - SEQUEDOCE: 14
          gagetiment trettataca aattatat
                                                                                     30
354 331: SE, 30 NO: .5
35. ....... LENGTH: 185
Nov. O. L. . THEE: DMA
dad 1.13 - ORGANISM: Badillus
THE HIGH REPUBLICE: 15
                                                                                     6.0
. . . . . .
          games: angu qeetgeaate gattgtttga gaaaagaaga agaccataaa aataccttgt
          rtigtostolig Gragggtatt tittatgotg topagactgt cogstgtgta aaaaatagga
                                                                                    110
11-3
. . ]
          athaandogis attigttatta tittactigat atgitaaaata taattigtat aagaaaatig
                                                                                    180
          ad 15d
                                                                                    135
344 4010 SEQ 10 NO: 16
.4: -1011 - LENGTH: 18:
2340 1.11 TYPE: DNA
347 -3113 - OFGANISM: Faillus
Jas - 3400 - SEQUENCE: 10-
                                                                                     6,0
.14
          squestiauqq asctqeaate qattqtttqa qaaaaqaaqa aqaccataaa aataccttgt
          ctateateng maggatatt tittatgeta tecagaetat cegetata aaaaaaaagaa
10
                                                                                    1.
: 1
                                                                                    1800
          ataanggggg yttgacatta ttttactgat atgtataata taatttgtat aagaaaatgg
          adata
                                                                                    14.5
1354 -0 10 - SEQ ID NO: 17
NET HULL FEMGTH: 185
DER ROTT FEE DNA
.in at 15, of GANISM: Facillus
. 15 K4000 SEQUENCE: 17
                                                                                     60
        - genetraling dectgeaate gattgtttgå gaaaagaaga agaccataaa aataccttgt
1.1
                                                                                    150
          cturcatoko wcagografi tittaigetg tecagacigi cegetgigia aaaaatagga
. . .
          atuaadeddd ditgaeatta tittactgat atgtataata taatiigtat aagaaaatgg
                                                                                    180
         agoto
                                                                                    165
164 C1C: SEQ II NO: 28
ies will length: 53
LOG - KLIKE TYPE: DNA
267 C 132 CkGANISM: Facillus
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/834,271A

DATE: 08/05/2003 TIME: 07:36:20

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt
Output Set: N:\CRF4\08052003\I834271A.raw

L:12 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD

STATISTICS SUMMARY

PATENT APPLICATION: US/09/834,271A

DATE: 08/05/2003 TIME: 07:36:20

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt
Output Set: N:\CRF4\08052003\I834271A.raw

Application Serial Number: US '09/834,271A

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 04-12-2001

Art Unit: OIPE

Software Application: FastSEQ3.0 Total Number of Sequences: 33 Total Nucleotides: 4542 Total Amino Acids: 0

Number of Errors: (Number of Warnings: 1 Number of Corrections: 0

MESSAGE SUMMARY

256 W: 1 (Invalid Numeric Header Field)